

STIC Biotechnology Systems Branch

CRF Problem Report

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) experienced a problem when processing the following computer readable form (CRF):

Application Serial Number: 10/502,510A

Filing Date: 7/23/04

Date Processed by STIC: 6/3/05

STIC Contact: Mark Spencer: Telephone: 571-272-2510; Fax: 571-273-0221

Nature of Problem:

The CRF (was):

- ☐ (circle one) Damaged or Unreadable (for Unreadable, see attached)
- ☐ Blank (no files on CRF) (see attached)
- ☐ Empty file (filename present, but no bytes in file) (see attached)
- ☐ Virus-infected. Virus name: _____ The STIC will not process the CRF.
- ☐ Not saved in ASCII text
- ☐ Sequence Listing was embedded in the file. According to Sequence Rules, submitted file should **only** be the Sequence Listing.
- ☒ Did not contain a Sequence Listing. (see attached sample)
- ☐ Other: _____

**PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM TO REDUCE ERRORS.
SEE BELOW FOR ADDRESS:**

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/efb/efs/downloads/documents.htm>>, EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05

(Sample of submitted file)



10/502,510A

- Pc T
- 8/9/04

CPProjectDataProject Data Screen: Current Application Number/Current Filing
Date: If Current Application Number or Current Filing Date are entered, both
must be entered. PMethods of screening molecular libraries and active molecules
identified thereby MBHB-04-585 (59.68.75763/001)
____-__-____3____yy____CProtein1Sequence____V3.3seq1
ArtificialIMETDTLLLWV LLLWVPGSTG D
21____ARNBCQEZGHILKMFPSTWYVX____PRT____yy____CCommentFe
ature____secretion signal
peptide____yy____COtherFeature____UNSUREsecretion signal
peptide211____V3.3seq2
ArtificiallAVGQDTQEVIVVPHSLPFKVVVISAILALVVTIISLIILIMLWQKKPR____ARNBCQEZGH
ILKMFPSTWYVX____PRT____E
____transmembrane domain____E
____UNSUREtransmembrane domain491____V3.3seq3
ArtificialMGWCT____ARNBCQEZGHILKMFPSTWYVX____PRT____E

p/s see error explanation
on next page.

Error Explanation: ✓

Please do not submit a PatentIn ".prj" file. It is a "work-in-progress" file only; you will need to submit a generated Sequence Listing. To generate one, go to "Application Steps" on PatentIn main menu. Click on "Generate Sequence Listing" and then click "Start" button. PatentIn will validate each sequence; hopefully, you will see "Successful completion" in the white box above the "Start" button.

You can then view the generated Sequence Listing by going to "Project" on main menu, and clicking on "View/Print Sequence Listing." A generated PatentIn Sequence Listing will have an ".ST25" file extension.

For further assistance, please contact Mark Spencer at 571-272-2510.